

PCT09

RAW SEQUENCE LISTING

DATE: 01/22/2002

PATENT APPLICATION: US/09/762,916

TIME: 13:22:57

Input Set : A:\92750-58.ST25.txt

Output Set: N:\CRF3\01182002\I762916.raw

ENTERED

3 <110> APPLICANT: Stichting voor de Technische Wetenschappen 5 <120> TITLE OF INVENTION: METHOD OF DETECTING A DNA SEQUENCE, A DNA SEQUENCE, A METHOD OF MAKING A DNA CONSTRUCT AND THE USE THEREOF 8 <130> FILE REFERENCE: 92750/58 10 <140> CURRENT APPLICATION NUMBER: US 09/762,916 12 <141> CURRENT FILING DATE: 1999-08-16 14 <150> PRIOR APPLICATION NUMBER: PCT/NL99/00518 16 <151> PRIOR FILING DATE: 1999-08-16 18 <150> PRIOR APPLICATION NUMBER: NL 1009862 20 <151> PRIOR FILING DATE: 1998-08-14 22 <160> NUMBER OF SEQ ID NOS: 5 24 <170> SOFTWARE: PatentIn version 3.0 27 <210> SEQ ID NO: 1 29 <211> LENGTH: 95 31 <212> TYPE: DNA 33 <213> ORGANISM: artificial sequence 35 <220> FEATURE: 37 <221> NAME/KEY: misc_feature 39 <222> LOCATION: (1)..(95) 41 <223> OTHER INFORMATION: synthetic sequence containing four binding sites for LexA from E. coli 44 <400> SEQUENCE: 1 46 gtcgactgct gtatataaaa ccagtggtta tatgtacagt acttgtactg tacatataac 60 95 48 cactggtttt atacagcaag cttggatccg tcgac 51 <210> SEQ ID NO: 2 53 <211> LENGTH: 73 55 <212> TYPE: DNA 57 <213> ORGANISM: artificial sequence 59 <220> FEATURE: 61 <221> NAME/KEY: primer_bind 63 <222> LOCATION: (1)..(73) 65 <223> OTHER INFORMATION: forward primer used to make human heat shock factor inducible promoter 68 <400> SEQUENCE: 2 70 aagcttggga gtcgaaactt ctggaatatt cccgaacttt cagccgacga cttataaaac 60 73 72 gccagggga agc 75 <210> SEQ ID NO: 3 77 <211> LENGTH: 76 79 <212> TYPE: DNA 81 <213> ORGANISM: artificial sequence 83 <220> FEATURE: 85 <221> NAME/KEY: primer_bind

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89 <223> OTHER INFORMATION: reverse primer used to make human heat shock factor inducible promoter 92 <400> SEQUENCE: 3 94 ccatggttta gcttccttag ctcctgaaaa tctcgccaag ctcccggggt ccgcgagaag 60 76 96 agctcggtcc ttccgg 99 <210> SEQ ID NO: 4 101 <211> LENGTH: 38 103 <212> TYPE: DNA 105 <213> ORGANISM: artificial sequence 107 <220> FEATURE: 109 <221> NAME/KEY: primer_bind 111 <222> LOCATION: (1)..(38) 113 <223> OTHER INFORMATION: forward PCR primer used to isolate DNA fragment from genomic Drosophila DNA 116 <400> SEQUENCE: 4 38 118 gatcaagctt atgatctgcg tatgatacca aatttctg 121 <210> SEQ ID NO: 5 123 <211> LENGTH: 36 125 <212> TYPE: DNA 127 <213> ORGANISM: artificial sequence 129 <220> FEATURE: 131 <221> NAME/KEY: primer_bind 133 <222> LOCATION: (1)..(36) 135 <223> OTHER INFORMATION: reverse PCR primer used to isolate DNA fragment from genomic 136 Drosophila DNA

36

138 <400> SEQUENCE: 5

140 gacaagetta cattgetggg egagetgege caateg

VERIFICATION SUMMARY

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